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See page 6-9



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**RAW SEQUENCE LISTING**

PATENT APPLICATION: US/10/081,327A

DATE: 07/25/2002

TIME: 08:42:12

Input Set : A:\SEQUENCE 081,327.txt

Output Set: N:\CRF3\07252002\J081327A.raw

3 <110> APPLICANT: Huang, Tim Hui-Ming  
5 <120> TITLE OF INVENTION: HIGH-THROUGHPUT METHODS FOR DETECTING DNA METHYLATION  
7 <130> FILE REFERENCE: 40629-2  
9 <140> CURRENT APPLICATION NUMBER: US 10/081,327A  
C--> 10 <141> CURRENT FILING DATE: 2002-07-11  
12 <150> PRIOR APPLICATION NUMBER: 09/497,855  
13 <151> PRIOR FILING DATE: 2000-02-04  
15 <150> PRIOR APPLICATION NUMBER: 60/120,592  
16 <151> PRIOR FILING DATE: 1999-02-18  
18 <150> PRIOR APPLICATION NUMBER: 60/118,760  
19 <151> PRIOR FILING DATE: 1999-02-05  
21 <160> NUMBER OF SEQ ID NOS: 54  
23 <170> SOFTWARE: PatentIn version 3.1  
25 <210> SEQ ID NO: 1  
26 <211> LENGTH: 630  
27 <212> TYPE: DNA  
28 <213> ORGANISM: Homo sapiens  
30 <220> FEATURE:  
31 <221> NAME/KEY: misc\_feature  
32 <222> LOCATION: (1)..(630)  
33 <223> OTHER INFORMATION: All n's are unknown  
36 <400> SEQUENCE: 1

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DEC 27 2002

**OFFICE OF PETITIONS**

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W--> 39	ataaaacaaa actttaccna tcgtatttgg gtaggggct ggggaatgga taactgttcc	120
W--> 41	ctctgctaaa atttttagcgt ntnaatcggg ggaatngagat ttgcataagt ttgacgggag	180
W--> 43	ggtcntgtnt tcgtttcaag tatgggtnat tgtccacacc ttttcttnaa gantngcgcg	240
W--> 45	gggggcctaa ttcgaanggt naanaaaana aggaagacng angtgccana atgnacccca	300
W--> 47	atccacanct tatggganan ctcccaacgc gttngatgca aacttgaatn tcnntactnt	360
W--> 49	cccctaaatn ncttgnggtt atcntggccn aactgtttcc tgnnttaaaa tnnttancg	420
W--> 51	ctccccattc cnetctannt acnaancggg aagcttnaat tttnaanncc nggggtncc	480
W--> 53	aatganttaa ctaaccccct tanttncctt nnnccccngc cegnttcnat ccggaaactn	540
W--> 55	ttttccccct ncnnttatta atccnccncc cccggaaaan gcggttgctt tntgggccct	600
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62	<212> TYPE: DNA	
63	<213> ORGANISM: Homo sapiens	
65	<220> FEATURE:	
66	<221> NAME/KEY: misc_feature	
67	<222> LOCATION: (1)..(913)	
68	<223> OTHER INFORMATION: All n's are unknown	
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      76 gctgggcgct ataatctgca gaacgtcccg ggccgaaaag gatggggaac acacctggag      180
W--> 78 ccagggcgaa gacatctaga gatgcaaagg gaatggaaga aaacccgngt aggttctctc      240
W--> 80 agagcaagac ccctttgctg gaaacatagg nttctaagcc cggtggtggg gagcaacagg      300
W--> 82 gtgggagaac tnaagactna gcctgaactt agcgctnccc cacttgctn cccatnaaga      360
W--> 84 tcccgtatatt ggcagggtcca gaccccgcca aacttcaaga ncccggcggt agggacaggc      420
W--> 86 aaacttgtag cgacangggg gaccggaacg gnacccccca ggaggggaan aaggggcngg      480
W--> 88 agatcagacc gttggtgnng gtngaattng gcnaaattnt tttnaacaca cntggtggng      540
W--> 90 tggaaccttn aatccctttt tagnctnct tgncttgac ctntnttagn ggggaaatcc      600
W--> 92 cctggggngn tntnntctt gcntngccaa ntggngggtg ngtnngnntt ttnttccatn      660
W--> 94 ntntnngggt tggcngatg tggggntntt tccngnttn ncngngnctt tttggggnnn      720
W--> 96 aanatngnct ttattntggg ggaaananaa agcttggnaa ttttttgggg aaanttgann      780
W--> 98 ccnnnngggt nggnaangnn ttntttnttg nngntctant ncgntnnggt ccttttttnc      840
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      107 <212> TYPE: DNA
      108 <213> ORGANISM: Homo sapiens
      110 <220> FEATURE:
      111 <221> NAME/KEY: misc_feature
      112 <222> LOCATION: (1)..(952)
      113 <223> OTHER INFORMATION: All n's are unknown
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W--> 121 gccatttcct tcctccgggg gtctttnta aaaganctcc ggnaccncct ccccatccc      180
W--> 123 tggagttgnt tctgcgcgga acttancctt ggggagagga ngtagcccg nctgtgacct      240
W--> 125 cggcctttcg ganngtgntc tgcgtntcct tactggcccc tccccttgct tccagatac      300
W--> 127 cctgtgtaac naactcgacc agnancggaa ggcgcggtta tgcattccagc anaaattgaa      360
W--> 129 aggtgcggaa ccggnaaaca aagatangag ggntnctgng tgccngccgt nctgcnactg      420
W--> 131 aatgaatgaa tagtgngact agtgaggaag nccaccatcc acngnggatc angatgtntg      480
W--> 133 anngataaaa ntggnaactg attttattgn gagagctccc aacgcattgg atgcatagct      540
W--> 135 tngagtatct ataagnnna cctaaataac ttgaccnaat catgggcaat aactggnncc      600
W--> 137 tgtngaaaaa tgnnatccct naccaanccn cacaacatnc taccnnaaa ccataaaagt      660
W--> 139 gtaaaaaaccn tgggggtccc caatnnantg aaccaacctc ncantnaatt nccntnnnc      720
W--> 141 ctcacnanc ccttttcnaa ccnaaaacc cttentaccc actncattaa atgaanccc      780
W--> 143 cccccccct aaaaanccn ntttncaaat tngnccccct tncnccntc ctntanntan      840
W--> 145 tnnactccna tanctcngnc ctatcntntt tcnnannaac nnnatcaanc tnanncaaaa      900
W--> 147 ncnaaaannc cnatteccc naaacccnan gaaaaccnna taaaaaacan tt          952
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      151 <211> LENGTH: 1059
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      153 <213> ORGANISM: Homo sapiens
      155 <220> FEATURE:
      156 <221> NAME/KEY: misc_feature
      157 <222> LOCATION: (1)..(1059)
      158 <223> OTHER INFORMATION: All n's are unknown
      161 <400> SEQUENCE: 4

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W--> 162 gnnnnnttga ancccttnag anccggnecc gacgtencat gctccccggcc gcctggccgc      60
164 gggatatcac tagtgccggc gcctgcaggt cgaccataat gagttgtagc aaatccacac      120
166 tttgtagttg ggaggaggag agctgggtgc caatgtttgt ccaactgttt atttttcgag      180
168 acccagccta ttgaggaagg cttgttcaag caggcttcgt tcctacaaac acacaaacat      240
170 ccttctgttt gcctacaaag aaaaggggac cagcgttga tttgcatgcg aatgaggagc      300
172 cccgggagga gacgtccttg cccatcaaag gccgcgtag cctggcggcc ctcagccttt      360
174 gttccagcac ctgctgcctt gcacggcctg acggggccgc gccccgcgt cctggctgcg      420
176 ccatccaggg tccacggctc actcgtccct cgctgcccc gcggccacac ctctgcagc      480
178 tcatacctca cccgccccgc ggccctgctt cccccagtct cctcctctcg ggagcaggcc      540
W--> 180 cgcagggtct gcaaggccaa gaacggaacc ccaaacgc an aggttgggga ggtgagccgg      600
182 caggacgcaa aactcggaat cgaggcggct tgggtttaca cgttccttta ctctggctca      660
184 cgctgggtctt cacctggtta tgggagagct tcccaacgcg ttggatgcat aacttgagta      720
W--> 186 ttctatagtg gcacctaaat aagctttggc gtaaacatgg gcatanctgg ntctctngnt      780
W--> 188 gaaattggta tccgnttcac aattccccac aacataacaa gcccggaagc nttaaagngg      840
W--> 190 aaaagctggg gtgcctaatt agggactnac tcacattaat tgcgntgggc ttacttgccc      900
W--> 192 nttttccaan cgggnaaaac ttctcngcca acttgcatat aatgaatcgg caaanccccc      960
W--> 194 ngnanaggcg nttnngnaa tgggcctttt ccttctcttg ttnnttaatn nntngngntg      1020
W--> 196 gccnttggtt nngcaacggg ntaantantn aaagggggg      1059
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200 <211> LENGTH: 895
201 <212> TYPE: DNA
202 <213> ORGANISM: Homo sapiens
204 <220> FEATURE:
205 <221> NAME/KEY: misc_feature
206 <222> LOCATION: (1)..(895)
207 <223> OTHER INFORMATION: All n's are unknown
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215 tctcaagtgt cctttggctc tggagctcct gggcccggtt tccgcatccg ggaggagaaa      180
W--> 217 ggagcgagcg cgcacggtgc ctcccagagg caggagcan aggagaggaa aggggaagac      240
W--> 219 tatgaaacgt nctaactaat gcgcgtactg aatgcgcatt cacgtggtca gttaccgatg      300
W--> 221 tgettctctc tctcctggcc cgcagggaca gacgcgccca gggacgtgcc agccccggnc      360
W--> 223 cgggctctgc acatttacta tgtgacctcc tgnagtcatt ctctctccca gtccatagatg      420
225 agatcttaga gtcacgcaa gcttgccctg tggaacagac taagtatctg actttgcggg      480
W--> 227 gatgtcctga tgcccacna aggacagaaa tcttgacac ttgcggatgn gctttatggg      540
W--> 229 agaagctcca acgcccgttg atgcatagct ttgggnattc tatagnacac cttnnatagct      600
W--> 231 ttggcgnaan tcatggncat ngntggnttn ctgngggaaa tggatatccg ttacaatttc      660
W--> 233 acacaacata caaccggaa gcantaaagt ngtaaagcct gggtgcccaa tnggggggng      720
W--> 235 ctaanttcca ttaattgggg tggcncannt gccgntttca gtcggggaaa acctggntgg      780
W--> 237 ccanctggnt tnttggnnnt ggccannccc ggggagaggn ggntttggga anttggngn      840
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244 <212> TYPE: DNA
245 <213> ORGANISM: Homo sapiens
247 <220> FEATURE:
248 <221> NAME/KEY: misc_feature
249 <222> LOCATION: (1)..(900)

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253 &lt;400&gt; SEQUENCE: 6

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258	actgaggcga	agaccgcgcc	ccggagtgtg	ggccgccttc	ccagtaaata	tgcttatgag	180
W--> 260	aggaagtgat	ggaccagatg	ctcaaggggt	agggactaga	ttttgcctta	canaggacaa	240
W--> 262	gtcagtctct	gaagagaagt	cgcggagagg	gacgcanaac	gcagtggggac	ggaanggtct	300
264	cggggaaagt	gggcttgccg	agaagtgcc	gggagctcgt	gggaagtgc	gagagccgct	360
266	gggattcggc	caggaaaaag	gagagtgcgc	gaacaagctt	cacagaaatg	gccttatggg	420
W--> 268	agaactccca	acgcgttgg	tgcntagctt	gantatctat	agtgtccctt	aaatacttgg	480
W--> 270	cgtaatcatg	gtcatanctg	tttctgtgt	taaatgttta	tccgctcnct	attccccgca	540
W--> 272	cctacgaacc	ngaaacctaa	ntntaaacct	ggggtgctaa	taattanctn	actcccatta	600
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W--> 276	gccccccccc	gaaaagggtt	gcttttnggg	ccttccnttc	ccccctaata	ctgncngcn	720
W--> 278	tcggtncggn	acgttccctc	ccaggngtat	tcgtttccga	acngggncct	gaaanctnac	780
W--> 280	aagcctnagg	cgacctnaag	gcntncngnt	tcnnggnccc	ccgananccc	aatnaccctt	840
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285 &lt;210&gt; SEQ ID NO: 7

286 &lt;211&gt; LENGTH: 730

287 &lt;212&gt; TYPE: DNA

288 &lt;213&gt; ORGANISM: Homo sapiens

290 &lt;220&gt; FEATURE:

291 &lt;221&gt; NAME/KEY: misc\_feature

292 &lt;222&gt; LOCATION: (1)..(730)

293 &lt;223&gt; OTHER INFORMATION: All n's are unknown

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W--> 301	gttgaggggtg	gtcccttggg	tcccagagaga	agctgcccct	cagaggctct	cananccan	180
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W--> 307	gagggagggtg	ccagaagtgtg	agccctgaag	ccctgctggc	ccctggggcg	aggcccanct	360
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W--> 315	aaccttttcc	cttttaaaat	ttttnccccc	cccttcccc	cccctnnaac	ccccanaana	600
W--> 317	nttttttccc	cnggnntncc	naaaatcanc	ccccattttt	tncccccccc	ccnctcngg	660
W--> 319	aaantttttc	tcntttttta	tntccccccg	gaaaaaagg	ttttntnccc	ccccccnaac	720
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324 &lt;210&gt; SEQ ID NO: 8

325 &lt;211&gt; LENGTH: 925

326 &lt;212&gt; TYPE: DNA

327 &lt;213&gt; ORGANISM: Homo sapiens

329 &lt;220&gt; FEATURE:

330 &lt;221&gt; NAME/KEY: misc\_feature

331 &lt;222&gt; LOCATION: (1)..(925)

332 &lt;223&gt; OTHER INFORMATION: All n's are unknown

335 &lt;400&gt; SEQUENCE: 8

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338 tcactagtgc ggccgcctgc aggtcgacca taatccctac accaaccctg ccaggagggt      120
W--> 340 tttctaangg cattttacag atgagcaaac tgtaccagcc agagtctcca aagcgcgggc      180
W--> 342 accccacact gntttnctnct tggcgngggg tgggggggtct cttagagggg gtaagagcac      240
W--> 344 caggagcccc aggcagcccc gntnccact ctggctgccc ggaccgtncg ggagccacgt      300
W--> 346 cccncccaa aggaggcccc ttacctgcgg gggcacctnc cgggcgcctc gcgtgtcggg      360
W--> 348 cccaaagcca ccgaggctgg gagecgtgg ctatactcgg tctctnctct gcccgnccgg      420
W--> 350 ccccgagctn ctactgcang cgggcccgtt gccntttctt ggttcnctt gctagncccc      480
W--> 352 cncngctgan ggccecatna ncmtncgg aggaggagg gaggcgctt ggtggccnga      540
W--> 354 ggcggcggtt ntattgctta cttgntcngg ggggtggggg nntctnagan tgggtcnanc      600
W--> 356 tnnntggggn ggcttnctnn gtngnncccn ctaangnggg gngcattnt tgcnnnngga      660
W--> 358 nnnngccccn nngnggggnt tnnnnnaaan gttttttgt ctgannnnca agcantngnn      720
W--> 360 tgganttant ttgngngggg angnnccnc nacnnggnnn gtannngttg nnnnagtttt      780
W--> 362 tnnnaggggc cnnanagnan ggnnngngnn nngntgatng ngnttgcnt ngntaggntn      840
W--> 364 gnnntnctnt tnnnatgnn nttnatnatn nnnngcnnna nntgngnagg gngatncntt      900
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369 <210> SEQ ID NO: 9
370 <211> LENGTH: 1047
371 <212> TYPE: DNA
372 <213> ORGANISM: Homo sapiens
374 <220> FEATURE:
375 <221> NAME/KEY: misc_feature
376 <222> LOCATION: (1)..(1047)
377 <223> OTHER INFORMATION: All n's are unknown
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383 gggccgcct gcaggtcgac cataactctc gaggtttcct cttcccccac ctcatctgc      120
385 tagccctgac gtgaggggat aggtgctctg tcgccgtatt cccatgtaga tacgacagat      180
387 ggaacgtaac tgcggttggc caggggctcc gcagtggctg gggcgccgga gtgcgtgct      240
389 ctgactttcc ggtgccgggc tctgtctcca ggctggaaat gtgctcctag ctctcctagg      300
391 gggacctgga cctaggggag gagattgggc ggagcccgcg gagtgtgccc aacctatgga      360
393 ccacgtggg cctggtctcc tgcgagcgt ccgggctgtc cctgccggtg catcggggtt      420
395 ccacgtcctg gggagagacg aagatgccct gggaaagata aacagccaag ttttgtcccc      480
397 ggaggaaaga agcttggaac tggtcctgt gtctgctgcg gggagggcca ggggataagt      540
399 gcgggtgctt gccgcctcc cttgtagaac ttggcgtctc gcacctttta taaacgccag      600
W--> 401 gatccgaggc cagaagttag ntggtggcca ggntagctgc tgaatgggtt gtggangaca      660
W--> 403 aggantgga tgctggccag ggaagatgtg cttgtcaggt angtaactcg gacaagaaga      720
W--> 405 actntggccc ctgaggaaga acagaacca aggcacttct ggagcaattt ctggcntggg      780
W--> 407 gtgataaaaag gaacttggtt tnaggcnaga cacatnctta tgggagaact tcccaacgcc      840
W--> 409 gttggatcat aactttgaga ttctataagg gccctaaaaa aantttggcg gaaannatgg      900
W--> 411 gcaaaaanttg ttcccggggg aaatngtann ccntttacaa attncncaa cntncgancc      960
W--> 413 cgaaccttaa anggnaaanc ccggggggcc taangngnga nctaactccc attaaatggn      1020
W--> 415 nnggcnnnat ngccntttt caanncg      1047
418 <210> SEQ ID NO: 10
419 <211> LENGTH: 634
420 <212> TYPE: DNA
421 <213> ORGANISM: Homo sapiens
423 <220> FEATURE:
424 <221> NAME/KEY: misc_feature
425 <222> LOCATION: (1)..(634)

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/081,327A

DATE: 07/25/2002  
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Input Set : A:\SEQUENCE 081,327.txt  
Output Set: N:\CRF3\07252002\J081327A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:2; N Pos. 17,48,78,102,109,229,271,312,318,337,350,356,401,436,452,470  
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Seq#:3; N Pos. 1,4,8,15,18,29,30,43,62,80,95,148,156,163,167,189,206,221  
Seq#:3; N Pos. 231,253,254,258,266,311,323,325,352,375,387,393,395,399,405  
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Seq#:3; N Pos. 557,558,577,597,598,605,610,613,614,621,627,630,639,647,670  
Seq#:3; N Pos. 685,686,688,701,704,706,711,714,716,717,719,726,728,738,743  
Seq#:3; N Pos. 744,755,764,776,778,796,800,801,805,812,814,822,825,828,833  
Seq#:3; N Pos. 836,837,840,842,843,849,853,857,859,866,868,873,874,876,877  
Seq#:3; N Pos. 881,882,883,889,892,894,895,901,903,908,909,912,913,921,928  
Seq#:3; N Pos. 930,939,940,950  
Seq#:4; N Pos. 2,3,4,5,6,12,18,22,27,37,580,766,771,778,795,831,838,868,885  
Seq#:4; N Pos. 901,910,915,926,955,961,963,965,971,976,978,1003,1004,1010  
Seq#:4; N Pos. 1011,1012,1014,1017,1019,1024,1031,1032,1041,1045,1048,1050  
Seq#:5; N Pos. 3,112,220,251,359,393,499,530,576,586,593,594,607,610,617  
Seq#:5; N Pos. 621,623,627,630,634,684,691,712,719,725,745,748,749,755,777  
Seq#:5; N Pos. 784,789,792,797,798,799,806,807,820,823,832,837,838,840,842  
Seq#:5; N Pos. 851,858,869,886,892,894  
Seq#:6; N Pos. 80,107,233,277,295,444,453,497,528,530,551,561,563,587,589  
Seq#:6; N Pos. 606,612,613,650,687,697,714,717,720,726,730,746,763,767,775  
Seq#:6; N Pos. 776,778,787,797,803,805,807,809,814,817,825,827,834,845,856  
Seq#:6; N Pos. 868,883,884,893,897,900  
Seq#:7; N Pos. 2,15,79,173,175,180,217,222,271,358,371,377,386,400,404,438  
Seq#:7; N Pos. 445,458,463,481,500,505,526,532,533,538,545,565,574,586,587  
Seq#:7; N Pos. 596,599,601,607,612,615,616,618,621,629,642,653,654,658,664  
Seq#:7; N Pos. 674,682,700,705,707,717,724  
Seq#:8; N Pos. 10,11,14,23,29,35,40,128,192,195,198,207,262,265,288,305,339  
Seq#:8; N Pos. 406,416,430,439,455,466,467,476,482,484,490,499,501,503,504  
Seq#:8; N Pos. 506,538,551,565,568,581,582,586,590,597,599,602,603,604,609  
Seq#:8; N Pos. 610,616,619,620,623,625,626,630,635,637,642,649,654,655,656  
Seq#:8; N Pos. 657,661,662,663,666,670,671,672,674,679,683,684,685,686,687  
Seq#:8; N Pos. 690,705,706,707,708,715,717,719,720,725,729,734,737,742,744

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/081,327ADATE: 07/25/2002  
TIME: 08:42:13Input Set : A:\SEQUENCE 081,327.txt  
Output Set: N:\CRF3\07252002\J081327A.raw

Seq#:8; N Pos. 745,749,751,754,755,758,759,760,764,765,767,771,772,773,774  
Seq#:8; N Pos. 782,783,784,792,793,795,798,800,803,804,805,807,809,810,811  
Seq#:8; N Pos. 812,814,819,821,823,828,829,831,833,838,840,842,843,844,846  
Seq#:8; N Pos. 849,852,853,854,855,859,860,861,863,864,867,870,871,872,873  
Seq#:8; N Pos. 877,878,879,881,882,885,887,892,896,898,901,903,905,908,909  
Seq#:8; N Pos. 913,915,917,919,924  
Seq#:9; N Pos. 6,9,23,621,633,656,665,702,724,776,802,807,816,883,895,896  
Seq#:9; N Pos. 907,925,929,930,933,944,947,952,954,958,972,975,979,994,996  
Seq#:9; N Pos. 998,1001,1020,1021,1022,1026,1027,1028,1031,1036,1044,1045  
Seq#:10; N Pos. 29,40,56,115,123,129,157,206,220,247,262,272,281,283,313  
Seq#:10; N Pos. 323,334,358,359,361,365,372,379,387,404,411,442,447,463,466  
Seq#:10; N Pos. 488,489,512,513,515,516,524,526,529,530,542,550,564,569,570  
Seq#:10; N Pos. 574,577,584,600,602,606,610,611,628,632  
Seq#:11; N Pos. 2,38,51,79,80,83,85,88,92,106,112,115,134,135,150,190,208  
Seq#:11; N Pos. 213,215,218,250,254,278,279,328,339,363,389,393,394,406,408  
Seq#:11; N Pos. 417,419,421,428,433,440,441,444,447,451,459,467,469,470,472  
Seq#:11; N Pos. 475,483,485,495,503,505,519,531,538,542,551,552,566,573,584  
Seq#:11; N Pos. 587,600,601,605,609,612,614,618,619,621,623,624,628,630,636  
Seq#:11; N Pos. 638  
Seq#:12; N Pos. 3,73,106,147,170,221,236,260,352,360,364,372,380,437,459  
Seq#:12; N Pos. 461,478,483,489,502,507,508,535,536,538,549,562,571,580,585  
Seq#:12; N Pos. 599,601,602,605,606,615,632,647,648,669,678,682,706,707,712  
Seq#:12; N Pos. 727,740,744,745,750,752,754,755,759,760,762,763,765,767,770  
Seq#:12; N Pos. 774,776,780,786,787,792,805,807,814,832,838,840,849,850,857  
Seq#:12; N Pos. 858,867,872,873,875,876,879,885,886,900,901,906,909,913,920  
Seq#:12; N Pos. 929,930,934,935,938,940,943,949,950,951,959,966,968,969,973  
Seq#:12; N Pos. 978,983,986,987,991,994,1003,1005,1006,1010,1017,1019,1020  
Seq#:12; N Pos. 1021,1024,1026,1027,1030,1033,1037,1041,1044,1045  
Seq#:13; N Pos. 132,147,173,188,190,196,198,213,235,261,268,293,297,312,325  
Seq#:13; N Pos. 337,340,353,400,418,424,430,431,444,446,476,479,481,518,522  
Seq#:13; N Pos. 539,542,555,569,572,576,579,600  
Seq#:14; N Pos. 33,74,121,145,165,170,179,182,185,197,203,206,226,238,256  
Seq#:14; N Pos. 271,279,299,300,310,317,326,331,343,347,367,377,381,387,390  
Seq#:14; N Pos. 410,415,417,418,423,435,438,440,443,450,451,455,460,478,487  
Seq#:14; N Pos. 489,491,494,511,512,513,515,519,522,523,528,534,537,539,540  
Seq#:14; N Pos. 544,549,552,555,559,561,563,569,584,587,588,596,599,600,602  
Seq#:14; N Pos. 608,609,611,617,619,625,632,635,636,640,651,655,657,664,665  
Seq#:14; N Pos. 675,680,685,687,689,690,693,694,708,713,719,723,724,730,732  
Seq#:14; N Pos. 736,737,740,742,745,746,748,750,754,758,760,767,773,774,777  
Seq#:14; N Pos. 778,780,781,783,787,791,795,800,803,811,812,813,818,824,825  
Seq#:14; N Pos. 832,834,836,839,844,846,848,851,858,859,860,863,869,870,871  
Seq#:14; N Pos. 872,873,877,880,884,885,895,896,901,902,904,909,910,913,915  
Seq#:14; N Pos. 916,919,920,923,924,927  
Seq#:15; N Pos. 5,202,212,242,247,254,255,256,275,292,305,307,319,332,336  
Seq#:15; N Pos. 341,384,385,398,420,429,447,448,451,452,458,464,472,485,488  
Seq#:15; N Pos. 490,503,506,516,528,533,542,544,550,559,563,567,575,584,598  
Seq#:15; N Pos. 600,608,609,612,617,621,624,625,630,633,649,656,657,658,663  
Seq#:15; N Pos. 665,666,673,677,678,684,696,697,712,715,716,717,719,729,731  
Seq#:15; N Pos. 734,738,741,742,743,744,770,771,787,788,789,790,792,794,795

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/081,327ADATE: 07/25/2002  
TIME: 08:42:13Input Set : A:\SEQUENCE 081,327.txt  
Output Set: N:\CRF3\07252002\J081327A.raw

Seq#:15; N Pos. 796,797,798,799,802,804,810,811,813,816,818,819,822,826,831  
Seq#:15; N Pos. 832,834,835,838,846,848,857,858,861,867,869,873,874,877,881  
Seq#:15; N Pos. 883  
Seq#:16; N Pos. 91,411,472,485,503,528,537,542,544,550,565,566,573,603,613  
Seq#:16; N Pos. 644,657,659,665,669,673,692,701  
Seq#:17; N Pos. 3,44,117,129,130,185,220,346,352,379,400,402,415,419,473  
Seq#:17; N Pos. 497,503,512,518,537,544,545,559,578,613,619,637,641,643,652  
Seq#:17; N Pos. 655,660,670,675,679,683,684,689,696,698,703,706,709,712,714  
Seq#:17; N Pos. 721,727,732,735,743,745,750,766,771,790,791,792,795,801,803  
Seq#:17; N Pos. 806,807,809  
Seq#:18; N Pos. 2,5,29,30,70,81,82,85,90,91,93,95,112,167,172,175,184,225  
Seq#:18; N Pos. 231,239,240,255,260,261,270,273,275,301,304,317,335,351,367  
Seq#:18; N Pos. 376,379,386,388,390,394,395,396,398,400,403,406,411,422,425  
Seq#:18; N Pos. 430,434,438,447,458,465,467,469,470,475,481,482,496,505,515  
Seq#:18; N Pos. 516,518,525,531,537,540,550,555,556,557,559,561,582,600,602  
Seq#:18; N Pos. 603,606,619,621,624  
Seq#:19; N Pos. 5,11,22,36,43,78,108,117,118,142,166,167,169,178,199,208  
Seq#:19; N Pos. 210,212,225,227,235,241,250,259,263,279,300,314,319,320,332  
Seq#:19; N Pos. 336,344,346,358,364,369,372,383,386,397,401,415,421,430,433  
Seq#:19; N Pos. 434,441,463,465,474,481,482,511,518,520,525,527,530,535,538  
Seq#:19; N Pos. 554,555,559,563,568,576,586,592,593,604,607,613,628,639,644  
Seq#:19; N Pos. 650  
Seq#:20; N Pos. 6,103,129,159,191,306,335,345,378,380,395,407,426,454,461  
Seq#:20; N Pos. 463,468,471,506,509,510,525,564,570,572,574,578,584,600,613  
Seq#:20; N Pos. 621,624,634,635,645,651,658,664,669,674,678,679,680  
Seq#:21; N Pos. 1,2,8,9,16,17,18,19,20,21,229,271,311,317,398,410,415,421  
Seq#:21; N Pos. 423,429,433,440,453,457,462,476,478,480,496,500,511,512,513  
Seq#:21; N Pos. 514,517,520,529,534,535,546,547,550,554,558,570,573,574,587  
Seq#:21; N Pos. 598,599,602,609,611  
Seq#:22; N Pos. 6,7,12,22,25,30,32,56,60,111,114,125,136,138,143,194,202  
Seq#:22; N Pos. 234,248,266,291,311,312,329,333,342,344,372,374,382,388,393  
Seq#:22; N Pos. 400,404,416,444,448,449,455,467,482,488,497,507,508,509,518  
Seq#:22; N Pos. 526,527,530,532,533,534,546,552,556,562,567,570,577,594,595  
Seq#:22; N Pos. 597,609,610,618,619,621,626,630,642,643,645,648,650,652,661  
Seq#:22; N Pos. 672,673,676,681,685,687,690,694,697,699,702,706,712,716,718  
Seq#:22; N Pos. 722,726,730,731,734,741,743,744,747,753,761,762,767,769,770  
Seq#:22; N Pos. 771,773,774,784,787,797,802,804,810,833,837,839,840,843,845  
Seq#:22; N Pos. 849,850,853,860,861,868,871,876,877,885,886,892,893,894,899  
Seq#:22; N Pos. 900,915,920,923,932,940,952,959,963,965,972,977,990,992,996  
Seq#:22; N Pos. 998,999,1004,1013,1017,1019,1021,1026,1029,1030,1034,1035  
Seq#:22; N Pos. 1039,1040,1058,1060,1061,1062,1064,1068,1074,1075,1077,1078  
Seq#:22; N Pos. 1080,1087,1088,1091,1095,1096,1101,1108,1116,1132,1141,1147  
Seq#:22; N Pos. 1149,1150,1157,1158,1164,1165  
Seq#:23; N Pos. 11,13,58,100,178,184,209,224,300,351,374,426,440,455,477  
Seq#:23; N Pos. 496,510,523,527,531,547,549,554,568,569,578,582,595,602,606  
Seq#:23; N Pos. 616,618,624  
Seq#:24; N Pos. 1,4,7,8,9,13,15,16,17,21,64,66,67,68,69,70,76,77,84,101,106  
Seq#:24; N Pos. 112,137,146,151,171,184,190,192,241,251,252,256,258,274,296  
Seq#:24; N Pos. 304,306,325,329,341,343,372,399,417,435,438,443,467,471,474



RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/081,327A

DATE: 07/25/2002  
TIME: 08:42:13

Input Set : A:\SEQUENCE 081,327.txt  
Output Set: N:\CRF3\07252002\J081327A.raw

Seq#:24; N Pos. 501,508,510,527,547,548,557,558,564,566,568,575,583,589,591  
Seq#:24; N Pos. 595,599,601,602,604,609,610,614,616,617,618,624,627,640,652  
Seq#:24; N Pos. 657,659,663  
Seq#:25; N Pos. 48,78,184,212,282,293,376,398,414,429,442,466,470,473,486  
Seq#:25; N Pos. 491,514,519,525,541,544,550,572,598,605,626  
Seq#:26; N Pos. 372,418,420,433,452,474,487,493,498,514,535,549,554,556,559  
Seq#:26; N Pos. 566,567,569,583,586  
Seq#:27; N Pos. 251,278,322,334,340,356,357,367,370,401,413,414,420,432,441  
Seq#:27; N Pos. 447,452,453,458,459,462,489,497,504,510,515,519,530,536,540  
Seq#:27; N Pos. 544,545,557,559,560,564,577,581,582,583,589,596,597  
Seq#:28; N Pos. 19,35,73,76,82,135,206,209,213,215,217,236,238,245,247,256  
Seq#:28; N Pos. 267,281,292,300,308,331,356,358,363,378,406,410,412,418,422  
Seq#:28; N Pos. 440,471,475,477,485,513,516,551,561,564,572,580,610  
Seq#:29; N Pos. 3,74,171,187,203,258,265,282,294,297,300,307,315,340,355  
Seq#:29; N Pos. 360,367,373,374,386,392,401,427,432,433,439,441,449,457,458  
Seq#:29; N Pos. 476,480,493,496,497,504,509,517,523,526,527,533,543,551,552  
Seq#:29; N Pos. 554,556,558,563,564,565,570,576,577,579,581,582  
Seq#:30; N Pos. 1,3,9,10,18,20,21,183,287,386,413,418,421,434,438,442,452  
Seq#:30; N Pos. 456,458,463,468,469,473,475,496,505,506,511,529,533,535,537  
Seq#:30; N Pos. 539,540,544,545,547,554,562,570,571,577,582,584,585,590,592  
Seq#:30; N Pos. 598,599,600,609,614,618,620  
Seq#:33; N Pos. 45,64,139,140,175,191,220,229,238,246,265,292  
Seq#:34; N Pos. 182  
Seq#:35; N Pos. 140,183  
Seq#:36; N Pos. 59,104,168,193,219,239,250,266,284  
Seq#:41; N Pos. 49,112,135,159

## VERIFICATION SUMMARY

PATENT APPLICATION: US/10/081,327A

DATE: 07/25/2002

TIME: 08:42:13

Input Set : A:\SEQUENCE 081,327.txt

Output Set: N:\CRF3\07252002\J081327A.raw

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:37 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:39 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:60  
L:41 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:120  
L:43 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:180  
L:45 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:240  
L:47 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:300  
L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:360  
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:420  
L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:480  
L:55 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:540  
L:57 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:600  
L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:60  
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:180  
L:80 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:240  
L:82 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:300  
L:84 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:360  
L:86 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:420  
L:88 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:480  
L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:540  
L:92 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:600  
L:94 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:660  
L:96 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:720  
L:98 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:780  
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:840  
L:102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:900  
L:117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:119 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:60  
L:121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:120  
L:123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:180  
L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:240  
L:127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:300  
L:129 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:360  
L:131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:420  
L:133 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:480  
L:135 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:540  
L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:600  
L:139 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:660  
L:141 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:720  
L:143 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:780  
L:145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:840  
L:147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:900  
L:162 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0  
L:180 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:540  
L:186 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:720  
L:188 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:780  
L:190 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:840

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/081,327A

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Input Set : A:\SEQUENCE 081,327.txt

Output Set: N:\CRF3\07252002\J081327A.raw

L:192 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:900  
L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:960  
L:196 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:1020